SEQUENCE LISTING

<110> VAN ELS, Cecile Antoinette Carola Maria VAN BLEEK, Gerarda Maria BOOG, Marie-Claire Josefa Pia

<120> MHC class II haplotype specific immunodominancy of peptides derived from RSV fusion (F) or attachment. (G) proteins

<130> 28902.0014

<140> (not yet assigned)

<141> 2004-12-17

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<151> 2002-06-20

<160> 29

<170> PatentIn version 3.1

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<213> respiratory syncytial virus

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Lys Glu Asn Lys Cys Asn Gly Thr Asp Ala Lys Val Lys Leu Ile Lys 65 . 70 75

Gln Glu Leu Asp Lys Tyr Lys Asn Ala Val Thr Glu Leu Gln Leu Leu . 85

Met Gln Ser Thr Pro Pro Thr Asn Asn Arg Ala Arg Arg Glu Leu Pro

Arg	Phe	Met 115	Asn	Tyr		Leu	Asn 120	Asn	Ala	Lys	Lys	Thr 125	Asn	Val	Thr
Leu	Ser 130	Lys	_	Arg		Arg 135	Arg	Phe	Leu	-	Phe 140	Leu	Leu	Gly	Val
Gly 145	Ser	Ala	Ile	Ala	Ser 150	Gly	Val	Ala	Val	Ser 155	Lys	Val	Leu	His	Leu 160
Glu	Gly	Glu	Val	Asn 165	_	Ile	Lys	Ser	Ala 170	Leu	Leu	Ser	Thr	Asn 175	Lys
Ala	Val	Val	Ser. 180	Leu	Ser.	Asn	Gly	Val 185	Ser	Val	Leu	Thr	Ser 190	Lys	Val
Leu	Asp	Leu 195	Lys		Tyr		Asp 200	_	Gln	Leu	Leu	Pro 205	Ile	Val	Asn
Lys.	Gln 210	Ser	Cys.	Ser	Ile	Ser 215	Asn	Ile	Glu	Thr	Val 220	Ile	Glu	Phe	Gln
Gln 225	Lys	Asn	Asn	Arg	Leu 230	Leu	Glu	Ile	Thr	Arg 235		Phe	Ser	Val	Asn 240
Ala	Gly	Val	Thr	Thr 245	Pro	Val	Ser	Thr	Tyr 250	Met	Leu	Thr.	Asn	Ser 255	Glu
Leu	Leu	Ser	Leu 260	Ile	Asn	Asp	Met	Pro 265	Ile			Asp	Gln 270	Lys	Lys
Leu	Met	Ser 275	Asn	Asn	Val.		Ile 280		Arg	Gln		Ser 285	Tyr	Ser	Ile
Met	Ser 290	Ile	Ile	Lys	Glu	Glu 295	Val	Leu	Ala	Tyr	Val 300	Val	Gln	Leu	Pro
Leu 305	Tyr	Gly	Val	Ile	Asp 310	Thr	Pro	Cys	Trp	Lys 315		His	Thr	Ser	Pro 320
Leu	Cys	Thr	Thr	Asn 325	Thr	Lys	Glu	Gly	Ser 330	Asn	Ile	Cys	Leu	Thr 335	Arg
Thr	Asp	Arg	Gly 340	Trp	Tyr	Cys	Asp	Asn 345	Ala	Gly	Ser	Val	Ser 350	Phe	Phe

Pro Gln Ala Glu Thr Cys Lys Val Gln Ser Asn Arg Val Phe Cys Asp Thr Met Asn Ser Leu Thr Leu Pro Ser Glu Ile Asn Leu Cys Asn Val 375 380 Asp Ile Phe Asn Pro Lys Tyr Asp Cys Lys Ile Met Thr Ser Lys Thr 390 395 Asp Val Ser Ser Val Ile Thr Ser Leu Gly Ala Ile Val Ser Cys 405 410 Tyr Gly Lys Thr Lys Cys Thr Ala Ser Asn Lys Asn Arg Gly Ile Ile 420 425 . 430 Lys Thr Phe Ser Asn Gly Cys Asp Tyr Val Ser Asn Lys Gly Met Asp 435 . 440 Thr Val Ser Val Gly Asn Thr Leu Tyr Tyr Val Asn Lys Gln Glu Gly 455 460 Lys Ser Leu Tyr Val Lys Gly Glu Pro Ile Ile Asn Phe Tyr Asp Pro 465 470 475 480 Leu Val Phe Pro Ser Asp Glu Phe Asp Ala Ser Ile Ser Gln Val Asn . . 490 495 Glu Lys Ile Asn Gln Ser Leu Ala Phe Ile Arg Lys Ser Asp Glu Leu Leu His Asn Val Asn Ala Gly Lys Ser Thr Thr Asn Ile Met Ile Thr 515 520 525 Thr Ile Ile Ile Val Ile Ile Val Ile Leu Leu Ser Leu Ile Ala Val 530 . 535 . . . 540 Gly Leu Leu Tyr Cys Lys Ala Arg Ser Thr Pro Val Thr Leu Ser 545 Lys Asp Gln Leu Ser Gly Ile Asn Asn Ile Ala Phe Ser Asn 565 570

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Leu	Asn	Leu 35	Lys	Ser	Val	Ala	Gln 40	Ile	Thr	Leu	Ser	4 =	Leu	Ala	Met
Ile	Ile 50	Ser	Thr	Ser	Leu	Ile 55	Ile	Ala	Ala	Ile		Phe		Ala	Ser
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Ser	Gln	Ile	Lys	Asn 85	Thr	Thr	Pro	Thr	Tyr 90	Leu	Thr		Asn		Gln
Leu	Gly	Ile	Ser 100	Pro	Ser	Asn	Pro	Ser 105	Glu	Ile	Thr	Ser	Gln 110	Ile	Thr
Thr	Ile	Leu 115	Ala	Ser	Thr	Thr	Pro 120	Gly	Val	_	Ser		Leu	Gln	Ser
Thr	Thr 130	Val	Lys	Thr	Lys	Asn 135	Thr	Thr	Thr	Thr	Gln 140		Gln	Pro	Ser
Lys 145	Pro	Thr	Thr	ГÀЗ	Gln 150	Arg	Gln	Asn	Lys	Pro 155	Pro	Ser	Lys	Pro	Asn 160
Asn	Asp	Phe	His	Phe 165	Glu	Val	Phe	Asn	Phe 170.	Val	Pro	Cys	Ser	Ile 175	Cys
Ser	Asn	Asn	Pro 180	Thr	Cys	Trp	Ala	Ile 185	Cys	Lys	Arg	Ile	Pro 190	Asn	Lys
Lys	Pro	Gly 195	Lys	Lys	Thr	Thr	Thr 200	Lys	Pro	Thr	Lys	Lys 205	Pro	Thr	Leu
Lys	Thr 210	Thr	Lys	Lys	Asp	Pro 215	Lys	Pro	Gln	Thr	Thr 220	Lys	Ser	Lys	Glu
Val 225	Pro	Thr	Thr	Lys	Pro 230	Thr	Glu	Glu	Pro	Thr 235	Ile	Asn	Thr	Thr	Lys 240

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Asn Pro Ser Pro Ser Gln Val Ser Thr Thr Ser Glu Tyr Pro Ser Gln
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